Atty. Dkt. No. 025098-0701 (Formerly 238/168)
Patent

IN THE CLAIMS

Please replace claims 1 and 49 with the following amended claims. A marked up version of the claims, indicating the changes made, is attached hereto as appendix A.

1. (Twice amended) A method for designing a specific polyamide

 $X_1 X_2 \ldots X_m \text{-} \gamma \text{-} X_{(m+1)} \ldots X_{(2m-1)} X_{2m} \text{-} R_1$

wherein

 $X_1, X_2, X_m, X_{(m+1)}, X_{(2m-1)}$, and X_{2m} are carboxamide residues forming carboxamide binding pairs $X_1/X_{2m}, X_2/X_{(2M-1)}, X_M/X_{M+1}$,

 γ is γ -aminobutyric acid or 2,4 diaminobutyric acid, and

R₁ is -NH(CH₂)₀₋₁₀₀NR₂R₃, -NH(CH₂)₀₋₁₂CONH(CH₂)₀₋₁₀₀NR₂R₃, or -NHR₂, where R₂ and R₃ are independently selected from the group consisting of H, Cl, NO, N-acetyl, benzyl, C₁₋₁₀₀ alkyl, C₁₋₁₀₀ alkylamine, C₁₋₁₀₀ alkyldiamine, C₁₋₁₀₀ alkylcarboxylate, C₁₋₁₀₀ alkenyl, a C₁₋₁₀₀ alkynyl, and C₁₋₁₀₀ alkyl-L, where L is selected from the group consisting of arylboronic acids, biotins, polyhistidines comprised from about 2 to 8 amino acids, haptens, solid phase supports, oligodeoxynucleotides, N-ethylnitrosourea, fluorescein, bromoacetamide, iodoacetamide, DL-α-lipoic acid, acridine, captothesin, pyrene, mitomycin, texas red, anthracene, anthrinilic acid, avidin, DAPI, and oligodeoxynucleotide, isosulfan blue, malachite green, psoralen, ethyl red, 4-(psoraen-8-yloxy)-butyrate, taartaric acid, and (+)-α-tocopheral, suitable for use as a DNA-binding ligand that is selective for identified target DNA-sequences 5'-WN₁N₂...N_mW-3' where m is an integer having a value from 3 to 6, the method comprising:

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- (a) identifying a target sequence of double stranded DNA having the form 5'-WN₁N₂ ... N_mW-3', N₁N₂ ... N_m being the sequence to be bound by carboxamide residues, wherein each N is independently chosen from the group A, G, C, and T, each W is independently chosen from the group A and T and m is an integer having a value from 3 to 6;
- (b) representing the identified sequence as 5'-Wab ... xW-3', wherein a is a first nucleotide to be bound by the X_1 carboxamide residue, b is a second nucleotide to be bound by the X_2 carboxamide residue, and x is the corresponding nucleotide to be bound by the X_m carboxamide residue;
- (c) defining a as A, G, C, or T to correspond to the first nucleotide to be bound by a carboxamide residue in the identified sequence;
- (d) selecting Im as the X_1 carboxamide residue and Py as the X_{2m} carboxamide residue if a = G;
- (e) selecting Py as the X_1 carboxamide residue and Im as the X_{2m} carboxamide residue if a = C;
- (f) selecting Hp as the X_1 carboxamide residue and Py as the X_{2m} carboxamide residue if a = T;
- (g) selecting Py as the X_1 carboxamide residue and Hz as the X_{2m} carboxamide residue if a = A; and
- (h) repeating steps c g for b through x until all carboxamide residues are selected; wherein Im is N-methylimidazole, Hp is, Py is N-methylpyrrole, A is adenine, G is guanine, C is cytosine, and T is thymine.



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Patent

49. (Amended)

A polyamide designed by the method of claim 1, having the structure:

FSUA)

$$R_4$$
 R_5
 R_5

 R_4 is selected from the group consisting of H, NH₂, SH, Cl, Br, F, N-acetyl, and N-formyl;

each R_5 is independently selected from the group consisting of H, $(CH_2)_{0-6}CH_3$, $(CH_2)_{0-6}CH_$

each R_6 is independently selected from the group consisting of H, NH_2 , OH, SH, Br, Cl, F, OMe, CH_2OH , CH_2SH , and CH_2NH_2 ;

 R_1 is $-NH(CH_2)_{0-100}NR_2R_3$, $-NH(CH_2)_{0-12}CONH(CH_2)_{0-100}NR_2R_3$, or $-NHR_2$, where R_2 and R_3 are independently selected from the group consisting of H, Cl, NO, N-acetyl, benzyl, C_{1-100} alkyl, C_{1-100} alkylamine, C_{1-100} alkyldiamine, C_{1-100} alkylcarboxylate, C_{1-100} alkenyl, a C_{1-100} alkynyl, and C_{1-100} alkyl-L, where L is selected from the group consisting of arylboronic acids, biotins, polyhistidines comprised from about 2 to 8 amino acids, haptens, solid phase supports, oligodeoxynucleotides, N-ethylnitrosourea, fluorescein, bromoacetamide, iodoacetamide, DL- α -